BISS Biodiversity Information Science and Standards

Biodiversity Information Science and Standards 2: e26080
doi: 10.3897/biss.2.26080

OPEN ACCESS

Conference Abstract

Automated Trait Extraction using ClearEarth, a Natural Language Processing System for Text Mining in Natural Sciences

Anne E Thessen[‡,§], Jenette Preciado[|], Payoj Jain[|], James H Martin[|], Martha Palmer[|], Riyaz Bhat[|]

‡ The Ronin Institute for Independent Scholarship, Monclair, NJ, United States of America
§ The Data Detektiv, Waltham, MA, United States of America
| University of Colorado Boulder, Boulder, CO, United States of America

Corresponding author: Anne E Thessen (annethessen@gmail.com)

Received: 22 Apr 2018 | Published: 22 May 2018

Citation: Thessen A, Preciado J, Jain P, Martin J, Palmer M, Bhat R (2018) Automated Trait Extraction using ClearEarth, a Natural Language Processing System for Text Mining in Natural Sciences. Biodiversity Information Science and Standards 2: e26080. https://doi.org/10.3897/biss.2.26080

Abstract

The cTAKES package (using the ClearTK Natural Language Processing toolkit Bethard et al. 2014, http://cleartk.github.io/cleartk/) has been successfully used to automatically read clinical notes in the medical field (Albright et al. 2013, Styler et al. 2014). It is used on a daily basis to automatically process clinical notes and extract relevant information by dozens of medical institutions. ClearEarth is a collaborative project that brings together computational linguistics and domain scientists to port Natural Language Processing (NLP) modules trained on the same types of linguistic annotation to the fields of geology, cryology, and ecology. The goal for ClearEarth in the ecology domain is the extraction of ecologically-relevant terms, including eco-phenotypic traits from text and the assignment of those traits to taxa. Four annotators used Anafora (an annotation software; https://github.com/weitechen/anafora) to mark seven entity types (biotic, aggregate, abiotic, locality, quality, unit, value) and six reciprocal property types (synonym of/has synonym, part of/has part, subtype/supertype) in 133 documents from primarily Encyclopedia of Life (EOL) and Wikipedia according to project guidelines (https://github.com/ClearEarthProject/AnnotationGuidelines). Inter-annotator agreement ranged from 43% to 90%. Performance of ClearEarth on identifying named entities in biology text overall was good (precision:

© Thessen A et al. This is an open access article distributed under the terms of the Creative Commons Attribution License (CC BY 4.0), which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

85.56%; recall: 71.57%). The named entities with the best performance were organisms and their parts/products (biotic entities - precision: 72.09%; recall: 54.17%) and systems and environments (aggregate entities - precision: 79.23%; recall: 75.34%). Terms and their relationships extracted by ClearEarth can be embedded in the new ecocore ontology after vetting (http://www.obofoundry.org/ontology/ecocore.html). This project enables use of advanced industry and research software within natural sciences for downstream operations such as data discovery, assessment, and analysis. In addition, ClearEarth uses the NLP results to generate domain-specific ontologies and other semantic resources.

Keywords

text mining, NLP, phenotype, ecology, ontology

Presenting author

Anne E Thessen

Presented at

TDWG 2018

Funding program

National Science Foundation: Data Infrastructure Building Blocks

Grant title

DIBBS: Porting Practical NLP and ML Semantics from Biomedicine to the Earth, Ice and Life Sciences

Hosting institution

University of Colorado Boulder

References

- Albright D, Lanfranchi A, Fredriksen A, Styler WF, Warner C, Hwang JD, Choi JD, Dligach D, Nielsen RD, Martin J, Ward W, Palmer M, Savova GK (2013) Towards comprehensive syntactic and semantic annotations of the clinical narrative. Journal of the American Medical Informatics Association 20 (5): 922-930. https://doi.org/10.1136/amiajnl-2012-001317
- Bethard S, Ogren P, Becker L (2014) ClearTK 2.0: Design Patterns for Machine Learning in UIMA. In LREC. LREC Int Conf Lang Resour Eval.
- Styler WI, Bethard S, Finan S, Palmer M, Pradhan S, de Groen P, Erickson B, Miller T, C L, Savova G, Pustejovsky J (2014) Temporal annotation in the clinical domain. Transactions of the Association for Computational Linguistics 2: 143-154.